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Amendments to the Claims

This listing of claims replaces all prior listings of claims in the application.

Listing of Claims:

- 15. (currently amended) A protein comprising two joined heterologous domains:
- a sequence non-specific double-stranded nucleic acid binding domain that comprises an amino acid sequence that has at least 75% 85% sequence identity to SEQ ID NO:2; and
 - a DNA polymerase domain

wherein the presence of the sequence non-specific double-stranded nucleic acid binding domain enhances the processivity of the polymerase domain compared to an identical protein that does not have the sequence non-specific double-stranded nucleic acid binding domain joined thereto.

- (previously presented) The protein of claim 15, wherein the sequence nonspecific double-stranded nucleic acid binding domain and the DNA polymerase domain are covalently linked.
 - 18.-21. (cancelled).
- (previously presented) The protein of claim 15, wherein the DNA polymerase domain has thermally stable polymerase activity.
- (previously presented) The protein of claim 15, wherein the DNA polymerase domain comprises a family A polymerase domain.

- 24. (previously presented) The protein of claim 23, wherein the family A polymerase domain is a *Thermus* polymerase domain.
- 25. (previously presented) The protein of claim 23, wherein the family A polymerase domain polymerase domain is a *Taq* polymerase domain.
- 26. (previously presented) The protein of claim 22, wherein the DNA polymerase domain is a ΔTaq domain.
- 27. (previously presented) The protein of claim 15, wherein the polymerase domain is a family B polymerase domain.
- 28. (previously presented) The protein of claim 27, wherein the family B polymerase domain is a *Pyrococcus* DNA polymerase I domain.
- (previously presented) The protein of claim 28, wherein the *Pyrococcus* polymerase domain is a *Pyrococcus furiosus* domain.
- 30. (currently amended) A protein comprising two joined heterologous domains:
- a sequence non-specific double-stranded nucleic acid binding domain that comprises an amino acid sequence that has at least 75% 85% sequence identity to the Sac7d sequence set forth in amino acids 7-71 of SEO ID NO:10; and
 - a DNA polymerase domain,
- wherein the presence of the sequence non-specific double-stranded nucleic acid binding domain enhances the processivity of the polymerase domain compared to an identical protein that does not have the sequence non-specific double-stranded nucleic acid binding domain joined thereto.

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- 31. (cancelled)
- (previously presented) The protein of claim 30, wherein the sequence nonspecific double-stranded nucleic acid binding domain and the DNA polymerase domain are covalently linked.

33. (cancelled)

- 34. (previously presented) The protein of claim 30, wherein the sequence nonspecific double-stranded nucleic acid binding domain comprises an amino acid sequence that has at least 90% sequence identity to the Sac 7d sequence set forth in SEQ ID NO:10.
- (previously presented) The protein of claim 30, wherein the DNA polymerase domain has thermally stable polymerase activity.
- (previously presented) The protein of claim 30, wherein the DNA polymerase domain comprises a family A polymerase domain.
- 37. (previously presented) The protein of claim 35, wherein the DNA polymerase domain is a *Thermus* polymerase domain.
- 38. (previously presented) The protein of claim 36, wherein the *Thermus* polymerase domain polymerase domain is a Taq polymerase domain.
- 39. (previously presented) The protein of claim 35, wherein the DNA polymerase domain is a ΔTaq domain.
- 40. (previously presented) The protein of claim 30, wherein the polymerase domain is a family B polymerase domain.

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- 41. (previously presented) The protein of claim 40, wherein the family B polymerase domain is a *Pyrococcus* DNA polymerase I domain.
- 42. (previously presented) The protein of claim 41, wherein the *Pyrococcus* polymerase domain is a *Pyrococcus furiosus* domain.